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Sequence Listing could not be accepted.

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Reviewer: Keisha Douglas

Timestamp: [year=2008; month=8; day=14; hr=9; min=55; sec=28; ms=876;]

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Reviewer Comments:

SEQUENCE LISTING

<110> APPLICANT: Ganymed Pharmaceuticals AG
Sahin, Ugur
Tureci, Oezlem
Koslowski, Michael

<120> TITLE OF INVENTION: Genetic Products Differentially Expressed In
Tumors And The Use Thereof

<130> FILE REFERENCE: 4883-0001

<140> CURRENT APPLICATION NUMBER:10537002

<141> CURRENT FILING DATE:2005-05-20

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globally throughout the sequence.

Application No: 10537002 Version No: 2.0

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Output Set:

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Total Errors: 0
No. of SeqIDs Defined: 141
Actual SeqID Count: 141

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SEQUENCE LISTING

<110> APPLICANT: Ganymed Pharmaceuticals AG
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<120> TITLE OF INVENTION: Genetic Products Differentially Expressed In Tumors And The Use Thereof

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<141> CURRENT FILING DATE:2005-05-20

<150> PRIOR APPLICATION NUMBER: PCT/EP2003/013091

<151> PRIOR FILING DATE: 2003-11-21

<150> PRIOR APPLICATION NUMBER: DE 102 54 601.0

<151> PRIOR FILING DATE: 2002-11-22

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<170> SOFTWARE: PatentIn version 3.1

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<213> ORGANISM: Homo sapiens

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<212> TYPE: DNA

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<210> SEO ID NO 7

<212> TYPE: DNA

<400> SEQUENCE: 7

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acagctgttt	tcaactacca	ggggctgtgg	cgctcctgtg	tccgagagag	ctctggcttc	180
accgagtgcc	ggggctactt	cacctgctg	gggctgccag	ccatgctgca	ggcagtgcga	240
gccctgatga	tcgtaggcat	cgtectgggt	gccattggcc	tcctggtatc	catctttgcc	300
ctgaaatgca	tccgcattgg	cagcatggag	gactctgcca	aagccaacat	gacactgacc	360
tccgggatca	tgttcattgt	ctcaggtctt	tgtgcaattg	ctggagtgtc	tgtgtttgcc	420
aacatgctgg	tgactaactt	ctggatgtcc	acagctaaca	tgtacaccgg	catgggtggg	480
atggtgcaga	ctgttcagac	caggtacaca	tttggtgcgg	ctctgttcgt	gggctgggtc	540
gctggaggcc	tcacactaat	tgggggtgtg	atgatgtgca	tcgectgccg	gggectggca	600
ccagaagaaa	ccaactacaa	agcggtttct	tatcatgcct	caggccacag	tgttgccctac	660
aagcctggag	gcttcaaggc	cagcactggc	tttggtcca	acacaaaaaa	caagaagata	720
tacgatggag	gtgccgcac	agaggacgag	gtacaatctt	atccttccaa	gcacgactat	780
ctgtaa						786

<210> SEO ID NO 8

<212> TYPE: DNA

<400> SEQUENCE: 8

tgcgcaccca	tggcgtgac	tgctgtcag	ggcttgggg	tctgtggttc	actgattggg	60
attgcgggca	tcattgtctgc	cacctgcatg	gaccagtggg	gcacccaaga	cttgtacaac	120
aaccccgtaa	cagctgtttt	caactaccag	gggctgtggc	gctcctgtgt	ccgagagagc	180

<210> SEQ ID NO 9

<212> TYPE: PRT

<400> SEQUENCE: 9

Met	Asn	Gly	Thr	Tyr	Asn	Thr	Cys	Gly	Ser	Ser	Asp	Leu	Thr	Trp	Pro
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Pro	Ala	Ile	Lys	Leu	Gly	Phe	Tyr	Ala	Tyr	Leu	Gly	Val	Leu	Leu	Val
			20					25					30		
Leu	Gly	Leu	Leu	Leu	Asn	Ser	Leu	Ala	Leu	Trp	Val	Phe	Cys	Cys	Arg
		35					40					45			
Met	Gln	Gln	Trp	Thr	Glu	Thr	Arg	Ile	Tyr	Met	Thr	Asn	Leu	Ala	Val
	50					55					60				
Ala	Asp	Leu	Cys	Leu	Leu	Cys	Thr	Leu	Pro	Phe	Val	Leu	His	Ser	Leu
65					70					75					80
Arg	Asp	Thr	Ser	Asp	Thr	Pro	Leu	Cys	Gln	Leu	Ser	Gln	Gly	Ile	Tyr
				85					90					95	
Leu	Thr	Asn	Arg	Tyr	Met	Ser	Ile	Ser	Leu	Val	Thr	Ala	Ile	Ala	Val
			100					105					110		
Asp	Arg	Tyr	Val	Ala	Val	Arg	His	Pro	Leu	Arg	Ala	Arg	Gly	Leu	Arg
		115					120					125			
Ser	Pro	Arg	Gln	Ala	Ala	Ala	Val	Cys	Ala	Val	Leu	Trp	Val	Leu	Val
	130					135					140				
Ile	Gly	Ser	Leu	Val	Ala	Arg	Trp	Leu	Leu	Gly	Ile	Gln	Glu	Gly	Gly
145					150					155					160

Phe Cys Phe Arg Ser Thr Arg His Asn Phe Asn Ser Met Arg Phe Pro
165 170 175
Leu Leu Gly Phe Tyr Leu Pro Leu Ala Val Val Val Phe Cys Ser Leu
180